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RAW SEQUENCE LISTING

DATE: 03/04/2002

PATENT APPLICATION: US/10/076,708

TIME: 16:06:07

Input Set : A:\Ep.txt

Output Set: N:\CRF3\03042002\J076708.raw

3 <110> APPLICANT: Sharma, Satish
 4 Rank, Kenneth
 6 <120> TITLE OF INVENTION: Assays for Accessing Aa-Tau Aggregation
 8 <130> FILE REFERENCE: 6322
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/076,708
 C--> 10 <141> CURRENT FILING DATE: 2002-02-15
 10 <160> NUMBER OF SEQ ID NOS: 7
 12 <170> SOFTWARE: PatentIn version 3.0
 14 <210> SEQ ID NO: 1
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42	ccaagggggc	tgatggtaaa	acgaagatcg	ccacaccgcg	gggagcagcc	cctccaggcc	720
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126 20 25 30
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129 35 40 45
131 Gln Thr Pro Thr Glu Asp Gly Ser Glu Glu Pro Gly Ser Glu Thr Ser
132 50 55 60
134 Asp Ala Lys Ser Thr Pro Thr Ala Glu Asp Val Thr Ala Pro Leu Val
135 65 70 75 80
137 Asp Glu Gly Ala Pro Gly Lys Gln Ala Ala Gln Pro His Thr Glu
138 85 90 95
140 Ile Pro Glu Gly Thr Thr Ala Glu Glu Ala Gly Ile Gly Asp Thr Pro
141 100 105 110
143 Ser Leu Glu Asp Glu Ala Ala Gly His Val Thr Gln Ala Arg Met Val
144 115 120 125
146 Ser Lys Ser Lys Asp Gly Thr Gly Ser Asp Asp Lys Lys Ala Lys Gly
147 130 135 140
149 Ala Asp Gly Lys Thr Lys Ile Ala Thr Pro Arg Gly Ala Ala Pro Pro
150 145 150 155 160
152 Gly Gln Lys Gly Gln Ala Asn Ala Thr Arg Ile Pro Ala Lys Thr Pro
153 165 170 175
155 Pro Ala Pro Lys Thr Pro Pro Ser Ser Gly Glu Pro Pro Lys Ser Gly
156 180 185 190
158 Asp Arg Ser Gly Tyr Ser Ser Pro Gly Ser Pro Gly Thr Pro Gly Ser
159 195 200 205
161 Arg Ser Arg Thr Pro Ser Leu Pro Thr Pro Pro Thr Arg Glu Pro Lys

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164 Lys Val Ala Val Val Arg Thr Pro Pro Lys Ser Pro Ser Ser Ala Lys
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167 Ser Arg Leu Gln Thr Ala Pro Val Pro Met Pro Asp Leu Lys Asn Val
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170 Lys Ser Lys Ile Gly Ser Thr Glu Asn Leu Lys His Gln Pro Gly Gly
171      260      265      270
173 Gly Lys Val Gln Ile Ile Asn Lys Lys Leu Asp Leu Ser Asn Val Gln
174      275      280      285
176 Ser Lys Cys Gly Ser Lys Asp Asn Ile Lys His Val Pro Gly Gly Gly
177      290      295      300
179 Ser Val Gln Ile Val Tyr Lys Pro Val Asp Leu Ser Lys Val Thr Ser
180 305      310      315      320
182 Lys Cys Gly Ser Leu Gly Asn Ile His His Lys Pro Gly Gly Gly Gln
183      325      330      335
185 Val Glu Val Lys Ser Glu Lys Leu Asp Phe Lys Asp Arg Val Gln Ser
186      340      345      350
188 Lys Ile Gly Ser Leu Asp Asn Ile Thr His Val Pro Gly Gly Gly Asn
189      355      360      365
191 Lys Lys Ile Glu Thr His Lys Leu Thr Phe Arg Glu Asn Ala Lys Ala
192      370      375      380
194 Lys Thr Asp His Gly Ala Glu Ile Val Tyr Lys Ser Pro Val Val Ser
195 385      390      395      400
197 Gly Asp Thr Ser Pro Arg His Leu Ser Asn Val Ser Ser Thr Gly Ser
198      405      410      415
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216 atgcaccaag accaagaggg tgacacggac gctggcctga aagctgaaga agcaggcatt      180
218 ggagacaccc ccagcctgga agacgaagct gctggtcacg tgaccaagc tcgcatggtc      240
220 agtaaaagca aagacgggac tggaagcgat gacaaaaaag ccaagggggc tgatggtaaa      300
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224 accaggattc cagcaaaaac cccgcccgtc ccaaagacac caccagctc tggatgaacct      420
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232 cccatgccag acctgaagaa tgtcaagtcc aagatcggct cactgagaa cctgaagcac      660
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236 tccaagtgtg gctcaaagga taatatcaaa caggtcccgg gaggcggcag tgtgcaaata      780
238 gtctacaaac cagttgacct gagcaaggtg acctccaagt gtggctcatt aggcaacatc      840
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246 ggggcggaga tcgtgtacaa gtgcacagtg gtgtctgggg acacgtctcc acggcatctc    1080
248 agcaatgtct cctccaccgg cagcatcgac atggtagact cgccccagct cgccacgcta    1140
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267 35 40 45
269 Gly Ile Gly Asp Thr Pro Ser Leu Glu Asp Glu Ala Ala Gly His Val
270 50 55 60
272 Thr Gln Ala Arg Met Val Ser Lys Ser Lys Asp Gly Thr Gly Ser Asp
273 65 70 75 80
275 Asp Lys Lys Ala Lys Gly Ala Asp Gly Lys Thr Lys Ile Ala Thr Pro
276 85 90 95
278 Arg Gly Ala Ala Pro Pro Gly Gln Lys Gly Gln Ala Asn Ala Thr Arg
279 100 105 110
281 Ile Pro Ala Lys Thr Pro Pro Ala Pro Lys Thr Pro Pro Ser Ser Gly
282 115 120 125
284 Glu Pro Pro Lys Ser Gly Asp Arg Ser Gly Tyr Ser Ser Pro Gly Ser
285 130 135 140
287 Pro Gly Thr Pro Gly Ser Arg Ser Arg Thr Pro Ser Leu Pro Thr Pro
288 145 150 155 160
290 Pro Thr Arg Glu Pro Lys Lys Val Ala Val Val Arg Thr Pro Pro Lys
291 165 170 175
293 Ser Pro Ser Ser Ala Lys Ser Arg Leu Gln Thr Ala Pro Val Pro Met
294 180 185 190
296 Pro Asp Leu Lys Asn Val Lys Ser Lys Ile Gly Ser Thr Glu Asn Leu
297 195 200 205
299 Lys His Gln Pro Gly Gly Gly Lys Val Gln Ile Ile Asn Lys Lys Leu
300 210 215 220
302 Asp Leu Ser Asn Val Gln Ser Lys Cys Gly Ser Lys Asp Asn Ile Lys
303 225 230 235 240
305 His Val Pro Gly Gly Gly Ser Val Gln Ile Val Tyr Lys Pro Val Asp
306 245 250 255
308 Leu Ser Lys Val Thr Ser Lys Cys Gly Ser Leu Gly Asn Ile His His
309 260 265 270
311 Lys Pro Gly Gly Gly Gln Val Glu Val Lys Ser Glu Lys Leu Asp Phe
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318 305          310          315          320
320 Arg Glu Asn Ala Lys Ala Lys Thr Asp His Gly Ala Glu Ile Val Tyr
321          325          330          335
323 Lys Ser Pro Val Val Ser Gly Asp Thr Ser Pro Arg His Leu Ser Asn
324          340          345          350
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327          355          360          365
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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date